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35	c	100..0	1455	4 US-09-170-470-10
36	c	100..0	1455	4 US-09-170-470-10
37	c	100..0	1455	4 US-09-193-068-10
38	c	100..0	1455	4 US-09-193-068-14
39	c	100..0	1455	4 US-09-183-412-19
40	c	100..0	1455	4 US-09-183-412-19
41	c	100..0	1455	4 US-09-354-191A-5
42	c	100..0	1455	4 US-09-290-734-10
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44	c	100..0	1689	1 US-07-991-8678-41
45	c	100..0	1689	2 US-08-544-332-41

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C	C	7	24	100.0	539	4
C	C	8	24	100.0	541	2
C	C	9	24	100.0	728	4
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C	C	11	24	100.0	993	1
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CG	CG	1	24	100.0	20	4
CG	CG	2	24	100.0	44	4
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CGT	CGT	8	24	100.0	541	2
CGT	CGT	9	24	100.0	728	4
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RESULT 1
RS-C9-657-481A-28/c
Sequence 28, Application US/09657481A
Patent No. 6,286,01
GENERAL INFORMATION:
APPLICANT: Brett P. Moria
APPLICANT: Lex M. Cowert
TITLE OF INVENTION: ANTISENSE MODULATION OF UBIQUITIN PROTEIN LIGASE (UPL)
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: PCT/US07/0087
CURRENT APPLICATION NUMBER: US/09/657,481A
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 93
SEQ ID NO: 28
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: OTHER INFORMATION: Antisense oligonucleotide
US-09657-481A-28

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c	2	24	100.0	44	4	78.99-291	874-3	Sequence 3, Appli
	3	24	100.0	56	4	US-09-291	874-15	Sequence 15, Appli
	4	24	100.0	539	4	US-09-605	885-310	Sequence 310, Appli
	5	24	100.0	539	4	US-09-439	913-310	Sequence 310, Appli
	6	24	100.0	539	4	US-09-352	616A-410	Sequence 310, Appli
	7	24	100.0	539	4	US-09-232	749A-310	Sequence 310, Appli
	8	24	100.0	591	2	US-08-342	766A-17	Sequence 17, Appli
	9	24	100.0	728	4	US-08-896	164-53	Sequence 53, Appli
	10	24	100.0	990	1	US-08-410	167A-1	Sequence 1, Appli
	11	24	100.0	993	1	US-08-705	337-1	Sequence 1, Appli
	12	24	100.0	993	1	US-08-705	337-2	Sequence 2, Appli

US 09 291 874 3
Sequence 3, Application US/09291874
; Patent No. 6436694

GENERAL INFORMATION:

APPLICANT: Francis P. Tally

; APPLICANT: Jianshi Tao

; APPLICANT: Xiaoya Shen

; APPLICANT: Jianwu Zhang

TITLE OF INVENTION: Gram-Positive Bacteria

FILE REFERENCE: C1198 03PMA2

CURRENT APPLICATION NUMBER: US/09/291,874

CURRENT FILING DATE: 1999-04-14

; EARLIER APPLICATION NUMBER: 60/122,949

; EARLIER FILING DATE: 1999-03-05

; EARLIER APPLICATION NUMBER: 09/227,687

; EARLIER FILING DATE: 1999-01-08

; EARLIER APPLICATION NUMBER: 60/107,751

; EARLIER FILING DATE: 1998-11-10

; EARLIER APPLICATION NUMBER: 60/101,718

; EARLIER FILING DATE: 1998-09-24

; EARLIER APPLICATION NUMBER: 60/100,211

; EARLIER FILING DATE: 1998-09-14

; EARLIER APPLICATION NUMBER: 60/094,698

; EARLIER FILING DATE: 1998-07-30

; EARLIER APPLICATION NUMBER: 60/089,828

; EARLIER FILING DATE: 1998-06-19

; EARLIER APPLICATION NUMBER: 60/085,844

; EARLIER FILING DATE: 1998-06-19

; EARLIER APPLICATION NUMBER: 60/085,844

; EARLIER FILING DATE: 1998-05-18

; EARLIER APPLICATION NUMBER: 60/084,698

; EARLIER FILING DATE: 1998-04-14

; EARLIER APPLICATION NUMBER: 60/076,638

; EARLIER FILING DATE: 1998-03-03

; EARLIER APPLICATION NUMBER: 60/070,965

; EARLIER FILING DATE: 1998-01-09

; SOFTWARE: FastSEQ for Windows Version 3.0

; SEQ ID NO: 15

; LENGTH: 56

Alignment Scores:

Pred. No.: 27.5

Score: 24.00

Length: 56

Matches: 5

Percent Similarity: 100.00%

; Conserv.:

; Pat. No.: 6,321,116

; Best Local Similarity: 100.00%

; Mismatches: 0

; Query Match: 100.00%

; Indels: 0

; Gaps: 0

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; RESULT: 4

; US-09-605-785-310

; Sequence: 310, Application US/09605785

; Patent No.: 6,321,116

GENERAL INFORMATION:

; APPLICANT: Xu, Jianchun

; APPLICANT: Dillon, Davin C.

; APPLICANT: Mitcham, Jennifer L.

; APPLICANT: Harzocker, Susan L.

; APPLICANT: Jiang, Yuqiu

; APPLICANT: Henderson, Robert A.

; APPLICANT: Kalos, Michael D.

; APPLICANT: Fanger, Gary R.

; APPLICANT: Reiter, Marc W.

; APPLICANT: Stolk, John A.

; APPLICANT: Day, Craig H.

; APPLICANT: Vedyick, Thomas S.

; APPLICANT: Carter, Derrick

; APPLICANT: Li, Samuel

; APPLICANT: Wang, Aljun

; APPLICANT: Skeky, Yasir A.W.

; APPLICANT: Beppler, William

TITLE OR INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

; FILE REFERENCE: 210121.427C16

; CURRENT APPLICATION NUMBER: US/09/291,874

; CURRENT FILING DATE: 2000-06-27

; EARLIER APPLICATION NUMBER: 60/122,949

; EARLIER FILING DATE: 1999-03-05

; EARLIER APPLICATION NUMBER: 60/094,698

; EARLIER FILING DATE: 1999-01-08

; EARLIER APPLICATION NUMBER: 60/089,828

; EARLIER FILING DATE: 1998-07-30

; EARLIER APPLICATION NUMBER: 60/085,844

; EARLIER FILING DATE: 1998-06-19

; EARLIER APPLICATION NUMBER: 60/100,211

; EARLIER FILING DATE: 1998-05-18

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; EARLIER FILING DATE: 1998-01-09

; NUMBER OF SEQ ID NOS: 28

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; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

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TYPE: DNA

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FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

; OTHER INFORMATION: oligonucleotide

; SEQ ID NO: 3

; LENGTH: 34

TYPE

NUMBER OF SEQ ID NOS: 835
 SOFTWARE: FastSEQ for Windows Version 3.0
 SEQ ID NO: 310
 LENGTH: 539
 TYPE: DNA
 ORGANISM: Homo sapien
 US-09-605-785-310

Alignment Scores:
 Pred. No.: 320 Length: 539
 Score: 24.00 Matches: 5
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: US-09-552-616A-310 Gaps: 0

US-09-856-070-18 (1-5) x US-09-605-785-310 (1-539)

Qy 1 LysGlugluLeuMet 5
 ; Sequence 310, Application US/09439313
 ; Patent No. 6329505

Db 464 AAGGAAGAACCTAAG 478

RESULT 5
 US-09-430-313-310

; Sequence 310, Application US/09439313

; GENERAL INFORMATION:

; APPLICANT: Xu, Jiatingchun
 ; APPLICANT: Dillon, Davin C.
 ; APPLICANT: Mitcham, Jennifer L.
 ; APPLICANT: Harlock, Susan Louise
 ; APPLICANT: Reed, Steven G.
 ; APPLICANT: Kalos, Michael
 ; APPLICANT: Fanger, Gary
 ; APPLICANT: Reiter, Mark
 ; APPLICANT: Soll, John

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
 ; TREATMENT OF PROSTATE CANCER

; FILE REFERENCE: 210121.427C9
 ; CURRENT APPLICATION NUMBER: US/09439313

; CURRENT FILING DATE: 1999-11-12
 ; NUMBER OF SEQ ID NOS: 575

; SOFTWARE: FastSEQ for Windows Version 3.0
 ; SEQ ID NO: 310

; LENGTH: 539

; TYPE: DNA

; ORGANISM: Homo sapien
 US-09-439-313-310

Alignment Scores:
 Pred. No.: 320 Length: 539
 Score: 24.00 Matches: 5
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 4

US-09-856-070-18 (1-5) x US-09-439-310 (1-539)

Qy 1 LysGlugluLeuMet 5
 ; Sequence 310, Application US/09439313

Db 464 AAGGAAGAACCTAAG 478

RESULT 6
 US-09-452-616A-310

; Sequence 310, Application US/09352616A

; GENERAL INFORMATION:

; APPLICANT: DILLON, DAVIN C.
 ; APPLICANT: HARLOCK, SUSAN LOUISE
 ; APPLICANT: JIANGQ, YUQI
 ; APPLICANT: MITCHAM, JENNIFER L.
 ; APPLICANT: REED, STEVEN G.
 ; APPLICANT: RITTER, MARK
 ; APPLICANT: SOLL, JOHN
 ; APPLICANT: SOLK, JOHN

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
 ; TREATMENT OF PROSTATE CANCER

; FILE REFERENCE: 210121.427C9
 ; CURRENT APPLICATION NUMBER: US/09352616A

; CURRENT FILING DATE: 1999-01-15
 ; NUMBER OF SEQ ID NOS: 326

; SOFTWARE: FastSEQ for Windows Version 3.0
 ; SEQ ID NO: 310

; LENGTH: 539

; TYPE: DNA

; ORGANISM: Homo sapien
 US-09-439-313-310

Alignment Scores:
 Pred. No.: 320 Length: 539
 Score: 24.00 Matches: 5
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 4 Gaps: 0

US-09-452-616A-310 (1-5) x US-09-452-616A-310 (1-539)

Qy 1 LysGlugluLeuMet 5
 ; Sequence 310, Application US/09352616A

Db 464 AAGGAAGAACCTAAG 478

RESULT 8
 US-09-432-766A-17

; Sequence 17, Application US/08332766A

; GENERAL INFORMATION:

; APPLICANT: JEFFREYS, ALEC J.
 ; APPLICANT: ARMOUR, JOHN
 ; TITLE OF INVENTION: SIMPLE TANDEM REPEATS
 ; NUMBER OF SEQUENCES: 125

APPLICATION NUMBER: US/08/896,164
 FILING DATE: JULY 17, 1997
 CLASSIFICATION: 424
 ATTORNEY/AGENT INFORMATION:
 NAME: Ng, 6218521man D. Hansson
 REFERENCE/DOCKET NUMBER: 100-5499 - JEL/NIH/SIH
 REGISTRATION NUMBER: 30,946
 TELEPHONE: (212) 688-9200
 TELEFAX: (212) 838-3884
 INFORMATION FOR SEQ ID NO: 53:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 728 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear
 US-08-896-164-53

Alignment Scores:
 pred. No.: 4.44 Length: 7
 score: 24.00 Matches: 5
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DH: 4 Gaps: 0

US-08-896-164-53 (1-53) x US-08-896-164-53 (1-728)

Qy 1 LysGluGluLeuMet 5
 ||||||| ||||| |||||
 Db 232 AAAGGAGAAATTAATG 246

RESULT 10
 US-08-410-167A.1
 Sequence 1, Application US/08410167A
 ; Patent No. 5773273

GENERAL INFORMATION:
 APPLICANT: Tokuto NISHINO, Shinichi OHNUMA, Manabu
 ; APPLICANT: Chikara OHNO, Chika ASADA, Yuka HIGI
 TITLE OF INVENTION: Geranylgeranyl-Diphosphate Synt
 ; Coding Thereror

NUMBER OF SEQUENCES: 4
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Kenyon & Kenyon
 STREET: One Broadway
 CITY: New York
 STATE: NY
 COUNTRY: US
 ZIP: 10004

COMPUTER READABLE FORM:
 MEDIUM TYPE: 3+ floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
 SOFTWARE: WordPerfect 6.1 Windows

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/410,167A
 FILING DATE: 24-MAR-1995
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 6-53804
 FILING DATE: 24-MAR-1994
 APPLICATION NUMBER: JP 6-315572
 FILING DATE: 25-NOV-1994
 ATTORNEY/AGENT INFORMATION:
 NAME: Edward W. Gresson
 REGISTRATION NUMBER: 18,918
 REFERENCE/DOCKET NUMBER: 918
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (212) 425-7200
 TELEFAX: (212) 415-5288
 INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 990 base pairs

TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear
 MOLECULE TYPE: DNA (genomic)
 HYPOTHETICAL: NO
 ORIGINAL SOURCE: NO
 ORGANISM: Sulfolobus acidocaldarius
 STRAIN: ATCC 33909

FEATURE: CDS
 NAME/KEY: CDS
 LOCATION: 1-990

Alignment Scores:
 Pred. No.: 619
 Score: 24.00
 Percent Similarity: 100.00%
 Best Local Similarity: 100.00%
 Query Match: 100.00%
 DB: 811 AAAGAGAGATAATG 825

US-09-856-070-18 (1-5) x US-09-410-167A-1 (1-940)

Qy 1 LysGluGluMet 5
 Db 811 AAAGAGAGATAATG 825

RESULT 11
 US-09-0705-377-1
 Sequence 1, Application US-09-0705-377-1
 Patent No. 5807725
 GENERAL INFORMATION:
 APPLICANT: OHIO, Chikara, ASADA, Chika, OHNUMA, Shinichi,
 APPLICANT: NISHINO, Tokuro, HIROKA, Kazuhiko, HEMMI, Hisashi
 TITLE OF INVENTION: Long-Chain Prenyl DiPhosphate Synthase
 NUMBER OF SEQUENCES: 9
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Kenyon & Kenyon
 STREET: 1025 Connecticut Avenue, N.W., Suite 600
 CITY: Washington
 STATE: DC
 COUNTRY: USA
 ZIP: 20036
 COMPUTER READABLE FORM:
 MEDIUM TYPE: 3.5 inch Diskette
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Wordperfect 6.1 Windows
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US-09-0705-377-1
 FILING DATE: 29-AUG-1996
 CLASSIFICATION: 435
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: JP 7247043
 FILING DATE: 01-SEP-1995
 ATTORNEY/AGENT INFORMATION:
 NAME: Toffonetti, Judith L.
 REGISTRATION NUMBER: 39,048
 REFERENCE/DOCKET NUMBER: 77670/442
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-429-1776
 TELEFAX: 202-429-0796
 INFORMATION FOR SEQ ID NO: 2:
 MOLECULE TYPE: Mutated genomic DNA
 LENGTH: 993
 SEQUENCE CHARACTERISTICS:
 TYPE: Nucleic acid
 STRANDEDNESS: Double
 TOPOLOGY: Linear
 ORIGINAL SOURCE: Sulfolobus acidocaldarius
 LENGTH: 993
 MATCHES: 5
 CONSERVATIVE: 0
 MISMATCHES: 0
 INDELS: 0
 GAPS: 0

Alignment Scores:
 Pred. No.: 621
 Score: 24.00
 Percent Similarity: 100.00%
 Best Local Similarity: 100.00%
 Query Match: 100.00%
 DB: 811 AAAGAGAGATAATG 825

US-09-856-070-18 (1-5) x US-09-0705-377-2 (1-933)

Qy 1 LysGluGluMet 5
 Db 811 AAAGAGAGATAATG 825

RESULT 13

US 08-705-377-3 Application US/08705377
 Sequence 3, Application US/08705377
 Patent No. 5807725

GENERAL INFORMATION:
 APPLICANT: OHIO, Chikara, ASADA, Chika, OHNUMA, Shinichi, NISHINO, Tokuzo, HIROKA, Kazutake, HEMMI, Hisashi
 TITLE OF INVENTION: Long-Chain Phosphatidyl Diphosphate Sythetase
 NUMBER OF SEQUENCES: 9
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Kenyon & Kenyon
 STREET: 1025 Connecticut Avenue, N.W., Suite 600
 CITY: Washington
 STATE: DC
 COUNTRY: USA
 ZIP: 20016

COMPUTER READABLE FORM:
 MEDIUM TYPE: 3.5 inch Diskette
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Wordperfect 6.1 Windows

CURRENT APPLICATION DATA:
 FILING DATE: 29-AUG-1996
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 7247043
 FILING DATE: 01-SEP-1995
 ATTORNEY/AGENT INFORMATION:
 NAME: Tolletetti, Judith L.
 REGISTRATION NUMBER: 39,048
 REFERENCE/DOCKET NUMBER: 77670/442
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-429-1776
 TELEFAX: 202-429-0796
 INFORMATION FOR SEQ ID NO: 4:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 993
 TYPE: Nucleic acid
 STRANDEDNESS: Double strand
 TOPOLOGY: Linear
 MOLECULE TYPE: Mutated genomic DNA
 US-08-705-377-4

Alignment Scores:
 Pred. No.: 621 Length: 994
 REFERENCE/DOCKET NUMBER: 77670/442 Matches: 5
 TELECOMMUNICATION INFORMATION: Percent Similarity: 100.00% Conservative: 0
 TELEPHONE: 202-429-1776 Best Local Similarity: 100.00% Mismatches: 0
 TELEFAX: 202-429-0796 Query Match: 100.00% Indels: 0
 INFORMATION FOR SEQ ID NO: 4: DB: 1 Gaps: 0
 SEQUENCE CHARACTERISTICS:
 LENGTH: 993
 TYPE: Nucleic acid
 STRANDEDNESS: Double strand
 TOPOLOGY: Linear
 MOLECULE TYPE: Mutated genomic DNA
 US-08-705-377-3

US-08-705-377-5

Alignment Scores:
 Pred. No.: 621 Length: 993
 Sequence 5, Application US/08705377
 REFERENCE/DOCKET NUMBER: 776725
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-429-1776
 TELEFAX: 202-429-0796
 INFORMATION FOR SEQ ID NO: 5:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 993
 TYPE: Nucleic acid
 STRANDEDNESS: Double strand
 TOPOLOGY: Linear
 MOLECULE TYPE: Mutated genomic DNA
 US-08-705-377-4

Alignment Scores:
 Pred. No.: 621 Length: 993
 Sequence 5, Application US/08705377
 REFERENCE/DOCKET NUMBER: 776725
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-429-1776
 TELEFAX: 202-429-0796
 INFORMATION FOR SEQ ID NO: 5:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 993
 TYPE: Nucleic acid
 STRANDEDNESS: Double strand
 TOPOLOGY: Linear
 MOLECULE TYPE: Mutated genomic DNA
 US-08-705-377-4

US 08-705-377-5 Application US/08705377
 Sequence 4, Application US/08705377
 Patent No. 5807725

GENERAL INFORMATION:
 APPLICANT: OHIO, Chikara, ASADA, Chika, OHNUMA, Shinichi, NISHINO, Tokuzo, HIROKA, Kazutake, HEMMI, Hisashi
 TITLE OF INVENTION: Long-Chain Phosphatidyl Diphosphate Sythetase
 NUMBER OF SEQUENCES: 9
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Kenyon & Kenyon
 STREET: 1025 Connecticut Avenue, N.W., Suite 600
 STATE: DC
 COUNTRY: USA
 ZIP: 20016

COMPUTER READABLE FORM:
 MEDIUM TYPE: 3.5 inch Diskette
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Wordperfect 6.1 Windows

CURRENT APPLICATION DATA:
 FILING DATE: 29-AUG-1996
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 7247043
 FILING DATE: 01-SEP-1995
 ATTORNEY/AGENT INFORMATION:
 NAME: Tolletetti, Judith L.
 REGISTRATION NUMBER: 39,048

REFERENCE/POCKET NUMBER: 77670/442
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-429-1776
TELEFAX: 202-429-0796
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 993
TYPE: Nucleic acid
STRANDEDNESS: Double strand
TOPOLOGY: linear
MOLECULE TYPE: Mutated genomic DNA
US-08-705-377-5

Alignment Scores:
pred. No.: 621 length: 993
Score: 24.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 1 Gaps: 0

US-09-856-070-18 (1-5) x US-08-705-377-5 (1-993)

QY 1 LysGluGluLeuMet 5
Db 811 AAAGAAGAAATTAATG 825

Search completed: January 16, 2003, 21:41:23
Job time: 19.1429 secs

